Reply to Restriction Requirement of May 6, 2003

## **Amendments to Specification**

## Please replac paragraph starting on pag 3, line 1:

Figure 1A-B depicts the amino acid sequence alignment between the N-Acetylglutamate Kinase from *Synechocystis sp.* (NCBI General Identifier No. 1652434; SEQ ID NO:11), the instant corn contig assembled of clones cr1n.pk0113.g3 and p0111.cipmf81r (SEQ ID NO:2), the instant rice clone rls72.pk0023.c6 (SEQ ID NO:4), the instant soybean clone sdp2c.pk010.h6 (SEQ ID NO:6) and the instant wheat contig assembled of clones wlk1.pk0019.d6 and wl1.pk0009.a7. The top row indicates with asterisks (\*) the amino acids conserved among all sequences and with plus signs (+) the amino acids conserved only among the plant sequences. Dashes are used by the program to maximize the alignment of the sequences. The bottom row contains the consensus sequence (SEQ ID NO:12) comprising all the amino acids which are conserved among all plant sequences.

Please replace Table 5, starting on page 18, line 19:

<u>TABLE 5</u>

Amino Acid Variability Among Plant N-Acetylglutamate Kinase

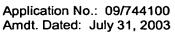
	Amino Acid Position in Consensus Sequence	Possible Amino Acids
	2	Leu, Met
	3	Leu, Ala
	4, <u>178<del>176</del></u>	Thr, Ala, Gly
,	<u>5</u>	Lys, none
	<u>6</u> 5	Pro, none
	<u>7</u> 6	<del>Trp</del> Tyr, His, none
	<u>8</u> 7	Leu, none
	9, <u>47</u> 8 <del>, 46</del>	Ser, Ala, none
	<u>10</u>	Asn, Ser, Ala, none
	<del>10</del> 11	Ser, Lys, Pro
	<del>11</del> <u>12</u>	<del>Lys</del> <u>Leu</u> , Ser, Thr, Ala

BI

Amino Acid Position in Consensus Sequence	Possible Amino Acids
<del>12</del> 13	Leu, Phe
<del></del>	Pro, Thr, none
14 <u>15</u>	Val, Leu, Asn
<del>15</del> 16	Pro, Leu
1 <del>6, 182</del> 17, 184	Ser, Cys
<del>17, 36<u>18, 37</u></del>	Pro, Thr, Ala
<del>18</del> <u>19</u>	Pro, Arg, Ser
<del>19</del> <u>20</u>	Pro, Val, Phe, Leu
<del>20, 30</del> <u>31</u>	Ser, Lys, Thr, Arg
<del>21</del> 22	Gly, Phe, Asn <u>, Ser</u>
<del>23, 50</del> <u>24, 51</u>	Thr, Ala, Asn
<del>2</del> 4 <u>25</u>	Leu, Pro, Lys
<del>25</del> <u>26</u>	Ser, Gly, Pro, Lys
<del>26</del> <u>27</u>	Ser, Pro, Gln, Ala
<del>27</del> <u>28</u>	Asn, Ala
<del>28</del> <u>29</u>	His, Gln, Arg
<del>29</del> <u>30</u>	Ala, Leu, Val
<del>31</del> <u>32</u>	Pro, Thr
<del>32</del> 33	Leu, IIe, Ser
<del>33</del> <u>34</u>	Ala, His
4 <del>, 49, 51, 52, 74, 194, 252, 265, 309</del>	Ala, Ser
35, 50, 52, 53, 75, 196, 254, 267, 311	
<del>35</del> <u>36</u>	Pro, Ser, Phe
<del>37</del> <u>38</u>	Cys, Ala, Ser, Pro
<del>38</del> <u>39</u>	Arg, Pro, Thr, His
<del>39</del> <u>40</u>	Arg, Gly, none
4 <del>0, 43<u>4</u>1, 44</del>	Arg, Leu
<u>4142</u>	Ser, Arg
4 <u>243</u>	Arg, Cys, His, Gly
44 <u>45</u>	Arg, Ala
4 <del>5, 253</del> 46, 2 <u>55</u>	lle, Leu, Val
47 <u>48</u>	Ala, Val, none
48 <u>49</u>	Thr, Val, none
<del>5</del> 3 <u>54</u>	Pro, Gln, Ser
<del>54, 57</del> <u>55, 58</u>	Ser, Ala, Pro

Amino Acid Position in	Descible Amine Aside
Consensus Sequence	Pro Ala Lau
<del>55</del> <u>56</u>	Pro, Ala, Leu
<del>56</del> <u>57</u>	Ser, Leu, Ala
<del>59</del> <u>60</u>	Ala, Glu, Gln
<del>60, 227, 428</del> <u>61, 229, 340</u>	Ala, Thr
<del>62</del> 63	Thr, Ser, none
<del>63, 191, 197, 269, 291</del>	Glu, Ala
<u>64, 193, 199, 271, 293</u>	
<del>64<u>65</u></del>	Ala, Ser, Gly
<del>65</del> <u>66</u>	Leu, Gln
<del>66</del> <u>67</u>	Ser, Tyr, Asn
<del>80, 82<u>81, 83</u></del>	Arg, Lys
86, 137, 154, 179, 226, 281	Val, lle
<u>87, 138, 156, 181, 228, 283</u>	
<del>106</del> <u>107</u>	Arg, Asn
<del>117, 215, 308</del> 118, 217, 310	Arg, His
<del>133</del> <u>134</u>	Leu, Gly, Gln
<del>135, 209</del> 136, 170	Val, <del>Ly</del> s <u>Leu</u>
<del>136</del> <u>137</u>	Gly, Asn
<del>138</del> <u>139</u>	Glu, Pro
<del>139</del> <u>141</u>	Gln, Val
<del>142, 210, 272</del> 144, 212, 274	Asp, Asn
<del>150</del> <u>152</u>	Leu, Asp
<del>151, 190</del> <u>153, 192</u>	Thr, Asn
<del>156</del> <u>158</u>	Glu, Ser
<del>161, 201</del> <u>163, 203</u>	Gly, Arg
<del>166</del> 168	Asn, Glu, Thr, Gln
<del>172</del> 174	Asn, Lys, Arg
<del>173</del> 175	lle, Leu, Lys, Pro
<del>174, 211, 237</del> 140, 176, 213, 239	Ala, Pro
<del>177, 228, 23</del> 4 <u>179, 230, 236</u>	Thr, Ser
<del>180, 221</del> <u>182, 223</u>	Gly, Ser
<del>18</del> 3 <u>185</u>	Gly, Trp, Arg
<del></del> <del>18</del> 4186	Lys, Met
<del>185, 292</del> 187, 294	Asp, Glu
<del>186, 230, 296</del> 188, 232, 298	Gly, Ala

BZ



_		
	Amino Acid Position in Consensus Sequence	Possible Amino Acids
-	189, 217, 249, 259, 261	lle, Leu
	<u>191, 219, 251,261, 263</u>	·
	<del>196</del> 198	Asn, Lys, Asp
	<del>198, 28</del> 4 <u>200, 286</u>	Lys, Ala
	<del>199</del> 201	Ala, Gly, Asp
	<del>202</del> 204	Phe, Tyr
B7-	<del>205</del> 207	Glu, Gly
	<del>207</del> 209	Ser, Trp, Ala, Thr
	<u>211</u>	<u>Val, Lys</u>
	<del>212, 220</del> 214, 222	Thr, Ala, Ser
	<del>216</del> 21, 218	Pro, Ser
	<del>219</del> 221	Ala, Asp
	<del>245</del> 247	Ala, Val
	<del>275</del> 277	Gly, Asp
	<del>280</del> 282	Val, Glu, Lys
	<del>288</del> <u>290</u>	Lys, Gln, Arg
	<del>290</del> 292	Val, Met
	<del>29</del> 4 <u>296</u>	Lys, Gln
	<del>30</del> 4 <u>306</u>	Glu, Asn, Gly
_	4 <del>05</del> 317	His, Ile

Please replace Figure 1 with the enclosed Figure 1A-B.

#### **Amended Claims**

## Please replace the Claims with the currently amended set:

Claim s 1-5 (currently canceled)

Claim 6.(currently amended) An isolated nucleic acid fragment polynucleotide comprising:

(a)a nucleotide sequence encoding a N-acetylglutamate kinase, wherein the polypeptide has an amino acid sequence of at least 95% sequence identity, based on the Clustal V method of alignment, when compared tohaving-the sequence set forth in SEQ ID NO:12, where:

Xaa at position 2 is Leu or Met,

Xaa at position 3 is Leu or Ala,

Xaa at position 4 or 476178 is Thr, Ala or Gly,

Xaa at position 5 is Lys or none,

Xaa at position 65 is Pro or none,

Xaa at position 76 is TyrTrp, His or none,

Xaa at position 87 is Leu or none,

Xaa at position 8, 9, or 4647 is Ser, Ala or none,

Xaa at position 10 is Asn, Ser, Ala, or none,

Xaa at position 1140 is Ser, Lys or Pro,

Xaa at position 1211 is LeuLys, Ser, Thr or Ala,

Xaa at position 1312 Leu or Phe,

Xaa at position <u>14</u>13 is Pro, Thr or none,

Xaa at position 1514 is Val, Leu or Asn,

Xaa at position 1615 is Pro or Leu,

Xaa at position 17 or 18416 or 182 is Ser or Cys,

Xaa at position 18 or 3717 or 36 is Pro, Thr or Ala,

Xaa at position 1918 is Pro, Arg or Ser,

Xaa at position 2019 is Pro, Val, Phe or Leu,

Xaa at position 3120 or 30 is Ser, Lys, Thr or Arg,

Xaa at position 2221 is Gly, Ser, Phe or Asn,

Xaa at position 24 or 51<del>23 or 50</del> is Thr, Ala or Asn,

Xaa at position 2524 is Leu, Pro or Lys,

Xaa at position 2625 is Ser, Gly, Pro or Lys,

Xaa at position 2726 is Ser, Pro, Gln or Ala,

Xaa at position 2827 is Asn or Ala,

Xaa at position 2928 is His, Gln or Arg,

Xaa at position 3029 is Ala, Leu or Val,

Xaa at position 3231 is Pro or Thr,

133

Xaa at position 3332 is Leu, Ile or Ser,

Xaa at position 3433 is Ala or His,

Xaa at position <u>35, 50, 52, 53, 75, 196, 254, 267, 311</u>34, 49, 51,

52, 74, 194, 252, 256 or 309 is Ala or Ser,

Xaa at position 3635 is Pro, Ser or Phe,

Xaa at position 3837 is Cys, Ala, Ser or Pro,

Xaa at position 3938 is Arg, Pro, Thr or His,

Xaa at position 4039 is Arg, Gly or none,

Xaa at position 41 or 4440 or 43 is Arg or Leu,

Xaa at position 4241 is Ser or Arg,

Xaa at position 4342 is Arg, Cys, His or Gly,

Xaa at position 4544 is Arg or Ala,

Xaa at position 46 or 25545 or 253 is Ile, Leu or Val,

Xaa at position 4847 is Ala, Val or none,

Xaa at position 4948 is Thr, Val or none,

Xaa at position 5453 is Pro, Gln or Ser,

Xaa at position 55 or 5854 or 57 is Ser, Ala or Pro,

Xaa at position 5655 is Pro, Ala or Leu,

Xaa at position 5756 is Ser, Leu or Ala,

Xaa at position 6059 is Ala, Glu or Gln,

Xaa at position 61, 229 or 340<del>60, 227 or 428</del> is Ala or Thr,

Xaa at position 6362 is Thr, Ser or none,

Xaa at position 64, 193, 199, 271, 293<del>63, 191, 197, 269 or 291</del>

is Glu or Ala,

Xaa at position 6564 is Ala, Ser or Gly,

Xaa at position 66 is 65 id Leu or Gln,

Xaa at position 6766 is Ser, Tyr or Asn,

Xaa at position 81 or 8380 or 82 is Arg or Lys,

Xaa at position <u>87, 138, 156, 181, 228, 28386, 137, 154, 179,</u>

#### 226, 281 is Val or Ile,

Xaa at position 107<del>106</del> is Arg or Asn,

Xaa at position 118, 217 or 310<del>117, 215 or 308</del> is Arg or His,

Xaa at position 134133 is Leu, Gly or Gln,

Xaa at position 136 or 170<del>135 or 209</del> is Val or LeuLys,

Xaa at position 137136 is Gly or Asn,

Xaa at position 139138 is Glu or Pro,

Xaa at position 141139 is Gln or Val,

Xaa at position <u>144, 212 or 274142, 210 or 272</u> is Asp or Asn,

B3

Xaa at position 152150 is Leu or Asp,

Xaa at position 153 or 192151 or 190 is Thr or Asn,

Xaa at position 158156 is Glu or Ser,

Xaa at position 163 or 203161 or 201 is Gly or Arg,

Xaa at position 168166 is Asn, Glu, Thr or Gln,

Xaa at position 174172 is Asn, Lys or Arg,

Xaa at position 175173 is Ile, Leu, Lys or Pro,

Xaa at position 140, 176, 213 or 239174, 211 or 237 is Ala or

Pro.

Xaa at position <u>179, 230 or 236 177, 228 or 234</u> is Thr or Ser,

Xaa at position 182 or 223180 or 221 is Gly or Ser,

Xaa at position 185183 is Gly, Trp or Arg,

Xaa at position 186184 is Lys or Met,

Xaa at position 187 or 294185 or 292 is Asp or Glu,

Xaa at position <u>188, 232 or 298 186, 230 or 296</u> is Gly or Ala,

Xaa at position 191, 219, 251, 261 or 263 189, 217, 249, 259 or

261 is lle or Leu.

Xaa at position 198196 is Asn, Lys or Asp,

Xaa at position 200 or 286198 or 284 is Lys or Ala,

Xaa at position 201199 is Ala, Gly or Asp,

Xaa at position 204<del>202</del> is Phe or Tyr,

Xaa at position 207205 is Glu or Gly,

Xaa at position 209<del>207</del> is Ser, Trp, Ala or Thr,

Xaa at position 211 is Val or Lys,

Xaa at position 214 or 222212 or 220 is Thr, Ala or Ser,

Xaa at position 21 or 218216 is Pro or Ser,

Xaa at position 221219 is Ala or Asp,

Xaa at position 247245 is Ala or Val,

Xaa at position 277275 is Gly or Asp,

Xaa at position 282280 is Val, Glu or Lys,

Xaa at position 290288 is Lys, Gln or Arg,

Xaa at position 292290 is Val or Met,

Xaa at position 296294 is Lys or Gln,

Xaa at position 306304 is Glu, Asn or Gly,

and Xaa at position 317405 is His or Ile.

Claim 7. (currently canceled)

Claim 8. (new) The polynucleotide of Claim 6 wherein:

Xaa at position 2 is Leu,

B3

Reply to Restriction Requirement of May 6, 2003

Xaa at position 3 is Leu, Xaa at position 4 is Thr, Xaa at position 5 is Lys, Xaa at position 6 is Pro, Xaa at position 7 is Tyr, Xaa at position 8 is Leu, Xaa at position 9 is Ser, Xaa at position 10 is Asn, Xaa at position 11 is Ser, Xaa at position 12 is Leu, Xaa at position 13 is Leu, Xaa at position 14 is Pro, Xaa at position 15 is Val, Xaa at position 16 is Pro, Xaa at position 17 is Ser, Xaa at position 18 is Pro, Xaa at position 19-is Pro, Xaa at position 20 is Pro, Xaa at position 21 is Ser, Xaa at position 22-is Gly, Xaa at position 24 is Thr, Xaa at position 25 is Leu, Xaa at position 26 is Ser, Xaa at position 27 is Ser, Xaa at position 28 is Asn, Xaa at position 29 is His, Xaa at position 30 is Ala, Xaa at position 31 is Ser, Xaa at position 32 is Pro, Xaa at position 33 is Leu, Xaa at position 34 is Ala, Xaa at position 35 is Ala, Xaa at position 36 is Pro, Xaa at position 37 is Thr, Xaa at position 38 is Cys, Xaa at position 39 is Arg, Xaa at position 40 is none, Xaa at position 41 is Arg,



Reply to Restriction Requirement of May 6, 2003

Xaa at position 42 is Ser, Xaa at position 43 is Arg, Xaa at position 44 is Leu, Xaa at position 45 is Arg, Xaa at position 46 is Ile, Xaa at position 47 is Ser, Xaa at position 48 is Ala, Xaa at position 49 is Thr, Xaa at position 50 is Ser, Xaa at position 51 is Thr, Xaa at position 52 is Ala, Xaa at position 53 is Ala, Xaa at position 54 is Pro, Xaa at position 55 is Ser, Xaa at position 56 is Pro, Xaa at position 57 is Ser Xaa at position 58 is Ser, Xaa at position 60 is Ala, Xaa at position 61 is Ala, Xaa at position 63 is Thr, Xaa at position 64 is Ala, Xaa at position 65 is Ser, Xaa at position 66 is Leu, Xaa at position 67 is Ser, Xaa at position 75 is Ala, Xaa at position 81 is Arg, Xaa at position 83 is Lys, Xaa at position 87 is Val, Xaa at position 107 is Arg, Xaa at position 118 is Arg, Xaa at position 134 is Leu, Xaa at position 136 is Val, Xaa at position 137 is Gly, Xaa at position 138 is Val, Xaa at position 139 is Glu, Xaa at position 140 is Pro, Xaa at position 141 is Gln, Xaa at position 144 is Asp,



Xaa at position 152 is Leu, Xaa at position 153 is Thr, Xaa at position 156 is Val, Xaa at position 158 is Glu, Xaa at position 163 is Gly, Xaa at position 168 is Asn, Xaa at position 170 is Val, Xaa at position 174 is Asn, Xaa at position 175 is Ile,

Xaa at position 211 is Val, Xaa at position 212 is Asp, Xaa at position 213 is Ala, Xaa at position 214 is Thr, Xaa at position 217 is His, Xaa at position 218 is Pro, Xaa at position 219 is Ile,

Xaa at position 176 is Ala, Xaa at position 178 is Gly, Xaa at position 179 is Thr, Xaa at position 181 is Ile, Xaa at position 182 is Gly, Xaa at position 184-is Cys, Xaa at position 185 is Gly, Xaa at position 186 is Lys, Xaa at position 187 is Asp, Xaa at position 188 is Ala, Xaa at position 191 is Ile, Xaa at position 192 is Thr, Xaa at position 193 is Ala, Xaa at position 196 is Ser, Xaa at position 198 is Asn, Xaa at position 199 is Ala, Xaa at position 200 is Ala, Xaa at position 201 is Ala, Xaa at position 203 is Gly, Xaa at position 204 is Phe, Xaa at position 207 is Glu, Xaa at position 209 is Ser,



Reply to Restriction Requirement of May 6, 2003

Xaa at position 221 is Ala, Xaa at position 222 is Ala, Xaa at position 223 is Gly, Xaa at position 228 is Ile, Xaa at position 229 is Ala, Xaa at position 230 is Thr, Xaa at position 232 is Ala, Xaa at position 236 is Thr, Xaa at position 239 is Ala, Xaa at position 247 is Ala, Xaa at position 251 is Ile, Xaa at position 254 is Ala, Xaa at position 255 is Val, Xaa at position 261 is Leu, Xaa at position 263 is Leu, Xaa at position 267 is Ser, Xaa at position 271 is Ala, Xaa at position 274 is Asn, Xaa at position 277 is Gly, Xaa at position 282 is Val, Xaa at position 283 is Val, Xaa at position 286 is Ala, Xaa at position 290 is Lys, Xaa at position 292 is Val, Xaa at position 293 is Ala, Xaa at position 294 is Asp, Xaa at position 296 is Lys, Xaa at position 298 is Ala, Xaa at position 306 is Glu, Xaa at position 310 is His, Xaa at position 311 is Ala, Xaa at position 317 is His, and Xaa at position 340 is Thr.

Claim 9. (new) The polynucleotide of Claim 6 wherein:

Xaa at position 2 is Leu, Xaa at position 3 is Leu, Xaa at position 4 is Ala,



Reply to Restriction Requirement of May 6, 2003

Xaa at position 5 is Lys, Xaa at position 6 is Pro, Xaa at position 7 is His, Xaa at position 8 is Leu, Xaa at position 9 is Ser, Xaa at position 10 is Ser, Xaa at position 11 is Ser, Xaa at position 12 is Ser, Xaa at position 13 Phe, Xaa at position 14 is none, Xaa at position 15 is Leu, Xaa at position 16 is Pro, Xaa at position 17 is Ser, Xaa at position 18 is Thr, Xaa at position 19-is Arg, Xaa at position 20 is Val, Xaa at position 21 is Ser, Xaa at position 22 is Ser, Xaa at position 24 is Ala, Xaa at position 25 is Pro, Xaa at position 26 is Gly, Xaa at position 27 is Pro, Xaa at position 28 is Asn, Xaa at position 29 is His, Xaa at position 30 is Ala, Xaa at position 31 is Lys, Xaa at position 32 is Pro, Xaa at position 33 is Ile, Xaa at position 34 is Ala, Xaa at position 35 is Ala, Xaa at position 36 is Ser, Xaa at position 37 is Pro, Xaa at position 38 is Ala, Xaa at position 39 is Pro, Xaa at position 40 is none, Xaa at position 41 is Arg, Xaa at position 42 is Arg, Xaa at position 43 is Cys,



Reply to Restriction Requirement of May 6, 2003

Xaa at position 44 is Leu, Xaa at position 45 is Arg, Xaa at position 46 is Leu, Xaa at position 47 is Ala, Xaa at position 48 is Val. Xaa at position 49 is Thr, Xaa at position 50 is Ser, Xaa at position 51 is Ala, Xaa at position 52 is Ala, Xaa at position 53 is Ala, Xaa at position 54 is Pro, Xaa at position 55 is Ala, Xaa at position 56 is Ala, Xaa at position 57 is Ser, Xaa at position 58 is Ser, Xaa at position 60 is Glu, Xaa at position 61 is Ala, Xaa at position 63 is none, Xaa at position 64 is Ala, Xaa at position 65 is Ala, Xaa at position 66 is Leu, Xaa at position 67 is Ser, Xaa at position 75 is Ala, Xaa at position 81 is Arg, Xaa at position 83 is Lys, Xaa at position 87 is Val, Xaa at position 107 is Arg, Xaa at position 118 is His, Xaa at position 134 is Leu, Xaa at position 136 is Val, Xaa at position 137 is Gly, Xaa at position 138 is Val, Xaa at position 139 is Glu, Xaa at position 140 is Pro, Xaa at position 141 is Gln, Xaa at position 144 is Asn, Xaa at position 152 is Leu, Xaa at position 153 is Asn,



Xaa at position 156 is Val, Xaa at position 158 is Glu, Xaa at position 163 is Arg, Xaa at position 168 is Glu, Xaa at position 170 is Leu, Xaa at position 174 is Lys, Xaa at position 175 is Leu, Xaa at position 176 is Pro, Xaa at position 178 is Gly, Xaa at position 179 is Ser, Xaa at position 181 is Val, Xaa at position 182 is Ser, Xaa at position 184-is Cys, Xaa at position 185 is Trp, Xaa at position 186 is Lys, Xaa at position 187 is Glu, Xaa at position 188 is Ala, Xaa at position 191 is Leu, Xaa at position 192 is Asn, Xaa at position 193 is Glu, Xaa at position 196 is Ser, Xaa at position 198 is Asn, Lys or Asp, Xaa at position 199 is Glu, Xaa at position 200 is Lys, Xaa at position 201 is Gly, Xaa at position 203 is Arg, Xaa at position 204 is Phe, Xaa at position 207 is Gly, Xaa at position 209 is Trp, Xaa at position 211 is Val, Xaa at position 212 is Asp, Xaa at position 213 is Ala, Xaa at position 214 is Thr, Xaa at position 217 is His, Xaa at position 218 is Pro, Xaa at position 219 is Ile, Xaa at position 221 is Ala,

Xaa at position 222 is Ala,

03

Reply to Restriction Requirement of May 6, 2003

Xaa at position 223 is Gly, Xaa at position 228 is Ile, Xaa at position 229 is Ala, Xaa at position 230 is Thr, Xaa at position 232 is Gly, Xaa at position 236 is Thr Xaa at position 239 is Ala, Xaa at position 247 is Ala, Xaa at position 251 is Ile, Xaa at position 254 is Ala, Xaa at position 255 is Val, Xaa at position 261 is Leu, Xaa at position 263 is Leu, Xaa at position 267 is Ser, Xaa at position 271 is Ala, Xaa at position 274 is Asn, Xaa at position 277 is Gly, Xaa at position 282 is Glu, Xaa at position 283 is Ile, Xaa at position 286 is Ala, Xaa at position 290 is Gln, Xaa at position 292 is Val, Xaa at position 293 is Ala, Xaa at position 294 is Asp, Xaa at position 296 is Gln, Xaa at position 298 is Ala, Xaa at position 306 is Glu, Xaa at position 310 is Arg, Xaa at position 311 is Ala, Xaa at position 317 is His, and

# Claim 10. (new) The polynucleotide of Claim 6 wherein:

Xaa at position 2 is Met, Xaa at position 3 is Ala, Xaa at position 4 is Gly, Xaa at position 5 is none, Xaa at position 6 is none,

Xaa at position 340 is Thr.



Reply to Restriction Requirement of May 6, 2003

Xaa at position 7 is none, Xaa at position 8 is none, Xaa at position 9 is Ala, Xaa at position 10 is Ala, Xaa at position 11 is Lys, Xaa at position 12 is Thr, Xaa at position 13n is Leu, Xaa at position 14 is Thr, Xaa at position 15 is Asn, Xaa at position 16 is Leu, Xaa at position 17 is Cys, Xaa at position 18 is Pro, Xaa at position 19-is Ser, Xaa at position 20 is Phe, Xaa at position 21 is Pro, Xaa at position 22 is Phe, Xaa at position 24 is Thr, Xaa at position 25 is Lys, Xaa at position 26 is Pro, Xaa at position 27 is Pro, Xaa at position 28 is Asn, Xaa at position 29 is Gln, Xaa at position 30 is Leu, Xaa at position 31 is Thr, Xaa at position 32 is Thr, Xaa at position 33 is Ser, Xaa at position 34 is His, Xaa at position 35 is Ala, Xaa at position 36 is Phe, Xaa at position 37 is Pro, Xaa at position 38 is Ser, Xaa at position 39 is Thr, Xaa at position 40 is Arg, Xaa at position 41 is Leu, Xaa at position 42 is Arg, Xaa at position 43 is His, Xaa at position 44 is Arg, Xaa at position 45 is Ala,



Xaa at position 46 is Ile, Xaa at position 47 is Ser, Xaa at position 48 is Ala, Xaa at position 49 is Val, Xaa at position 50 is Ala, Xaa at position 51 is Asn, Xaa at position 52 is Ala, Xaa at position 53 is Ala, Xaa at position 54 is Gln, Xaa at position 55 is Pro, Xaa at position 56 is Pro, Xaa at position 57 is Leu, Xaa at position 58 is Ala, Xaa at position 60 is Ala, Xaa at position 61 is Thr, Xaa at position 63 is Thr, Xaa at position 64 is Glu, Xaa at position 65 is Gly, Xaa at position 66 is Gln, Xaa at position 67 is Tyr, Xaa at position 75 is Ser, Xaa at position 81 is Lys, Xaa at position 83 is Arg, Xaa at position 87 is Ile, Xaa at position 107 is Asn, Xaa at position 118 is Arg, Xaa at position 134 is Gly, Xaa at position 136 is Leu, Xaa at position 137 is Asn, Xaa at position 138 is Ile, Xaa at position 139 is Pro, Xaa at position 140 is Ala, Xaa at position 141 is Val, Xaa at position 144 is Asp, Xaa at position 152 is Asp, Xaa at position 153 is Thr, Xaa at position 156 is Ile, Xaa at position 158 is Ser,



Reply to Restriction Requirement of May 6, 2003

Xaa at position 163 is Gly, Xaa at position 168 is Thr, Xaa at position 170 is Val, Xaa at position 174 is Asn, Xaa at position 175 is Lys, Xaa at position 176 is Ala, Xaa at position 178 is Ala, Xaa at position 179 is Thr, Xaa at position 181 is Val, Xaa at position 182 is Gly, Xaa at position 184-is Ser, Xaa at position 185 is Gly, Xaa at position 186 is Met, Xaa at position 187 is Asp, Xaa at position 188 is Gly, Xaa at position 191 is Leu, Xaa at position 192 is Thr, Xaa at position 193 is Ala, Xaa at position 196 is Ala, Xaa at position 198 is Lys, Xaa at position 199 is Ala, Xaa at position 200 is Ala, Xaa at position 201 is Asp, Xaa at position 203 is Gly, Xaa at position 204 is Tyr, Xaa at position 207 is Glu, Xaa at position 209 is Ala, Xaa at position 211 is Val, Xaa at position 212 is Asp, Xaa at position 213 is Pro, Xaa at position 214 is Ala, Xaa at position 217 is Arg, Xaa at position 218 is Ser, Xaa at position 219 is Leu, Xaa at position 221 is Asp, Xaa at position 222 is Thr, Xaa at position 223 is Ser, Xaa at position 228 is Val,



Reply to Restriction Requirement of May 6, 2003

Xaa at position 229 is Thr, Xaa at position 230 is Ser, Xaa at position 232 is Ala, Xaa at position 236 is Ser, Xaa at position 239 is Pro, Xaa at position 247 is Val, Xaa at position 251 is Leu, Xaa at position 254 is Ser, Xaa at position 255 is Leu, Xaa at position 261 is Ile, Xaa at position 263 is Leu, Xaa at position 267 is Ala, Xaa at position 271 is Glu, Xaa at position 274 is Asn, Xaa at position 277 is Asp, Xaa at position 282 is Lys, Xaa at position 283 is Ile, Xaa at position 286 is Lys, Xaa at position 290 is Lys, Xaa at position 292 is Met, Xaa at position 293 is Glu, Xaa at position 294 is Asp, Xaa at position 296 is Lys, Xaa at position 298 is Gly, Xaa at position 306 is Asn, Xaa at position 310 is Arg, Xaa at position 311 is Ser, Xaa at position 317 is Ile, and Xaa at position 340 is Ala.

# Claim 11. (new) The polynucleotide of Claim 6 wherein:

Xaa at position 2 is Leu, Xaa at position 3 is Leu, Xaa at position 4 is Thr, Xaa at position 5 is Lys, Xaa at position 6 is Pro, Xaa at position 7 is His, Xaa at position 8 is none,



Xaa at position 9 is none, Xaa at position 10 is none, Xaa at position 11 is Pro, Xaa at position 12 is Ala, Xaa at position 13 is Leu, Xaa at position 14 is Thr, Xaa at position 15 is Leu, Xaa at position 16 is Pro, Xaa at position 17 is Ser, Xaa at position 18 is Ala, Xaa at position 19 is Ser, Xaa at position 20 is Leu, Xaa at position 21 is Pro, Xaa at position 22 is Phe, Xaa at position 24 is Asn, Xaa at position 25 is Leu, Xaa at position 26 is Lys, Xaa at position 27 is Ala, Xaa at position 28 is Ala, Xaa at position 29 is Arg, Xaa at position 30 is Val, Xaa at position 31 is Arg, Xaa at position 32 is Pro, Xaa at position 33 is Leu, Xaa at position 34 is Ala, Xaa at position 35 is Ser, Xaa at position 36 is Ser, Xaa at position 37 is Ala, Xaa at position 38 is Pro, Xaa at position 39 is His, Xaa at position 40 is Gly, Xaa at position 41 is Arg, Xaa at position 42 is Arg, Xaa at position 43 is Gly, Xaa at position 44 is Leu, Xaa at position 45 is Arg, Xaa at position 46 is Val, Xaa at position 47 is none,

133

Reply to Restriction Requirement of May 6, 2003

Xaa at position 48 is none, Xaa at position 49 is none, Xaa at position 50 is Ser, Xaa at position 51 is Ala, Xaa at position 52 is Ser, Xaa at position 53 is Ser, Xaa at position 54 is Ser, Xaa at position 55 is Ser, Xaa at position 56 is Leu, Xaa at position 57 is Ala, Xaa at position 58 is Pro, Xaa at position 60 is Gln, Xaa at position 61 is Ala, Xaa at position 63 is Ser, Xaa at position 64 is Ala, Xaa at position 65 is Ala, Xaa at position 66 is Leu, Xaa at position 67 is Asn, Xaa at position 75 is Ser, Xaa at position 81 is Lys, Xaa at position 83 is Arg, Xaa at position 87 is Ile, Xaa at position 107 is Arg, Xaa at position 118 is Arg, Xaa at position 134 is Gln, Xaa at position 136 is Val, Xaa at position 137 is Gly, Xaa at position 138 is Val, Xaa at position 139 is Glu or Pro, Xaa at position 140 is Pro, Xaa at position 141 is Gln, Xaa at position 144 is Asn, Xaa at position 152 is Leu or Asp, Xaa at position 153 is Thr or Asn, Xaa at position 156 is Val or Ile, Xaa at position 158 is Glu or Ser, Xaa at position 163 is Gly or Arg, Xaa at position 168 is Gln,



Reply to Restriction Requirement of May 6, 2003

Xaa at position 170 is Leu, Xaa at position 174 is Arg, Xaa at position 175 is Pro, Xaa at position 176 is Ala, Xaa at position 178 is Thr, Xaa at position 179 is Thr, Xaa at position 181 is Val, Xaa at position 182 is Gly, Xaa at position 184-is Cys, Xaa at position 185 is Arg, Xaa at position 186 is Lys, Xaa at position 187 is Asp, Xaa at position 188 is Gly, Position 190 is Ile, Xaa at position 191 is Leu, Xaa at position 192 is Thr, Xaa at position 193 is Glu, Xaa at position 196 is Ser, Xaa at position 198 is Asp, Xaa at position 199 is Ala, Xaa at position 200 is Ala, Xaa at position 201 is Ala, Xaa at position 203 is Gly, Xaa at position 204 is Phe, Xaa at position 207 is Glu, Xaa at position 209 is Thr, Xaa at position 211 is Lys, Xaa at position 212 is Asn, Xaa at position 213 is Pro, Xaa at position 214 is Ser, Xaa at position 217 is His, Xaa at position 218 is Pro, Xaa at position 219 is Ile, Xaa at position 221 is Ala, Xaa at position 222 is Ser, Xaa at position 223 is Ser, Xaa at position 228 is Ile, Xaa at position 229 is Ala,



Xaa at position 230 is Thr, Xaa at position 232 is Ala, Xaa at position 236 is Thr, Xaa at position 239 is Ala, Xaa at position 247 is Ala, Xaa at position 251 is Ile, Xaa at position 254 is Ala, Xaa at position 255 is Ile, Xaa at position 261 is Leu, Xaa at position 263 is Ile, Xaa at position 267 is Ser, Xaa at position 271 is Ala, Xaa at position 274 is Asp, Xaa at position 277 is Gly, Xaa at position 282 is Glu, Xaa at position 283 is Ile, Xaa at position 286 is Ala, Xaa at position 290 is Arg, Xaa at position 292 is Val, Xaa at position 293 is Ala, Xaa at position 294 is Glu, Xaa at position 296 is Lys, Xaa at position 298 is Gly, Xaa at position 306 is Gly, Xaa at position 310 is Arg, Xaa at position 311 is Ala, Xaa at position 317 is His, and

Xaa at position 340 is Thr.

B3